R. Mitim



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/498,556C

DATE: 04/15/2003 TIME: 13:22:09

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Cutput Set: N:\CRF4\04152003\I498556C.raw

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? <1105 APPLICANT: Corvas International , Inc.
              Vlasuk, George Phillip
      Ġ
                                                             · (* f* -11
      ţ,
              Stanssens, Fatrick Eric Hugo
             Messens, Joris Hila Lieven
      ۴.
             - Lauwereys, Marc Josef
             Laroche, Yves Rene
             Jespers, Laurent Stephane
             Gansemans, Yannick Georges Jozef
     10
     1.
            - Mowile, Matthew
     1...
             Bergur, Peter W.
     14 -1200 TITLE OF INVENTION: MEMATODE-EMTRACTED SERINE PROTEASE INHIBITORS AND
ANTICOAGULANT PROTEIN
     16 ×1300 FILE REFERENCE: 018813/0272487
     18 -: 140: CUFFERM APPLICATION NUMBER: 09/498,556C
     13 - 1410 CUFFENT FILING DATE: 2005-04-01
     11 -150 PRIOR APPLICATION NUMBER: 09/809,455
     17 - 180. PRIOR FILING DATE: 1997-04-17.
     04 - 150 - PETOF APPLICATION NUMBER: PCT/US95/13231
     IS HIST. PRIOR FILLING DATE: 1995-10-17
     17 -1150 - PRIOR APPOLIMATION NUMBER: 08/486,399
                                                               ENTERED
     te RISI - PRIOR FILING CATE: 1995-06-05
     so -150 - PRIOR APPLICATION NUMBER: 06/456,397
     31 -151 FFIOF FILING SATE: 1995-06-05
     33 - 150 - FFIOF APPLICATION NUMBER: 08/465,380
     34 - 181 - FALOR FILING DATE: 1995-06-05
     -8 0150 - PRIOR APPRICATION NUMBER: 09/461,965
     -7 -: 151 - FAIOF FILSH'S DATE: 1995-06-06
     39 - 150 - PETOF APPLICATION NUMBER: 08/326,110
     49 - 0181 - FRIOR BILLING DATE: 1994-10-18
     4. HIGH - NUMBER OF SEQ ID NOS: 357
     44 <170 - SOFTWARE: Patentin version 3.1
     47 K21 + SEO ID NO: 1
     48 RUH1 - LENGTH: 234
     49 MAIL - TYPE: DNA
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     13 HOG - SENUEDCE: 1
     94 wagginatano oggaqtijtgg tgagaatgia tggotogaog actgtggaac tcagaagoca 60
     ні бустыддемы ngigonutya gyaacoochi gaggaggaag atoogatatg cegeteaegt 120
     no qythattta: naodtostgo tigogtatjo aaagaoggat totacagaga caeggigato 180
     ый түргүстэгд төүрүндүй agaatgogad baabatgaga ttatabatgt otga. 234
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     64 HILL: LENGTH: 228
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ยย์ <215> ORGANISM: Andyclostoma caninum

65 SC1. > TYPE: DNA

TIME: 13:22:09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/498,556C

Input Set: A:\Corvas-sequence listing.txt
Output Set: N:\CRF4\04152003\I498556C.raw

68 <400> SEQUENCE: 2 70 aaggeatado oggaqtiqtgg tgagaatgaa tggotogaog totgtggaac taagaaqooa 60 71 typogagyona agtycaytga qyaayagyag gaagatooga tatgoogato attitottyt $120\,$ 74 occupatocog otgottoogt atgogaagad ggattotada gagadaeggt gateggegad 180 76 tq-q-haayq aayaagaatg cgaccaabat gagattatac atgtotga -79 (210) NEQ ID NO: 3 80 -0011: LENGTH: 461 81 HOLLEY TYPE: DNA 81 - (1) 0 - OF CANISM: Andyclostoma caninum 84 - CLL OF FEATURE: 85 -CTOUR MAME/REY: COR 86 ACADA LOCATION: (..)..(821) SH RULLUR FRATURE: 8 + KLII - NAME/REY: MOD RED 96 KILL FORTHER INFORMATION: ADAMARS CDNA sequence 90 -0400 - BEQUENCE: 3 94 quarticognit actain was a sty aug atg off tab get atc get ata atg 51 Met Lys Met Leu Tyr Ala Ile Ala Ile Met GL. 97 tit, the city gia the tite type ago goa aga aca gity agg aag goa tao 99 98 Pho heu Leu Val Sor Leu Cys Ser Ala Arg Thr Val Arg Lys Ala Tyr 20 100 day day tot ggt dag aut gaa tog oto gad gad tot gga act dag aag 147 101 Pro Glu Cys Gly Glu Arn Glu Trp Leu Asp Asp Cys Gly Thr Gln Lys 30 10... 10, occurred gain god and the antique gain coe cot gain gain gain at cog 195 104 Pro Cys Gla Ala Lys Cys Asn Gla Gla Pro Pro Gla Gla Gla Asp Pro 50 5.5 100 ata the eye toa cut gut the tha tha bot got the gta the ada 243 107 Ile Cys Ard Ser Arg Gly Cys Leu Leu Pro Pro Ala Cys Val Cys Lys 10s vo 65 10% que oquitto tao aga que aby gtg ato ggb que tgt gtt agg gua jua 291 110 Asp Gly Fhe Tyr Arg Asp Thr Val Ile Gly Asp Cys Val Arg Glu Glu 111 75 11.0 qualityo gad caa but gug att ata dat gtd tgaabgagaaa gbaabaataabd 344 11% Glu Cys Asp Gln His Glu Ile Ile His Val 1:: 1.) 11% assysticca action metringeasasted staginggar greteriting egreeyasts 404 110 grittragitty atgittingta agaabtboty otggagagaa taaagottto baabtoo 114 K 10 - SEQ ID NO: 4 1.00 -0.111 - LENGTH: 77 101 -0010 - TYPE: PRP 1.1 Ull3 - ORGANISM: Aleyelostoma caninum 104 <400 - SEQUENCE: 4 136 Lys Ala Tyr Pro 310 Cys Sly Glu Asn Glu Trp Leu Asp Asp 127 1 1) 135 Cys (ly Thr Gln Lys Pro Cys Glu Ala Lys Cys Asn Glu Glu 21.1 1.11 15 130 Pro Pro Glu Glu Glu Aup Pro Ile Cys Ard Ser Arg Gly Cys

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Input Set : A:\Corvas-sequence listing.txt
Output Set: N:\CRF4\04152003\I498556C.raw

35 13.1 100 Leu beu Pro Pro Ala Cys Val Cys Lys Asp Gly Phe Tyr Arg 5.0 133 45 134 Asp Thr Val Ile Gly Asp Cys Val Arg Glu Glu Glu Cys Asp 135 60 65 130 Glr. His Glu The Ile His Val 137 7 r, 140 -010: SEQ ID NO: 5 141 -CL11: LENGTE: 455 140 -00120 TYPE: DNA 14: -0:12: ORGANISM: Andyclostoma caninum $145 - 1... \odot \cdot$ FEATURE: 146 FILL TO NAME/FEY: CDS 147 - LOGATION: (11)..(315) 149 CHION FEATURE: 150 - U.T. NAME/HEY: MOD RES 181 - C. W. OTHER INFORMATION: AdaMAP6 cDNA sequence 155 - 400 - SEQUENCE: 5 150 graftcoget amactosac a anglasg anglett tad get and get ata ang 51 Mot Lys Met Leu Tyr Ala Ile Ala Ile Met 156157 1^{4} % (if one only gray too tha ingo ago aca aga aca gity agg aag gos tac 99100 Phot Leu Leu Val Ser Leu Cys Ser Thr Arg Thr Val Arg Lys Ala Tyr .21) 161 coq qay tgt ggt gar aat gaa tyg oto gac gto tgt gga act aag aag 147 162 Pro-Gli Cys Gly Glu Ash Gli Trp Leu Asp Val Cys Gly Thr Lys Lys 3.0 4.0 35 164 das tigo gag god ang tigo agti gag gaa gag gag gaa gat oog ata tigo 195 165 Pro dys 3lu Ala Lys Cys Ser Giu Glu Glu Glu Glu Asp Pro Ile Cys 4.5 5 C $167~\mathrm{dyr}$ that that that being gift bed got got the gita the galace gas gift 243168 And Ber Phe Ser Cys Pro Gly Pro Ala Ala Cys Val Cys Glu Asp Gly 169 60 65 170 tip tao aga gab aog gig ato ggo gao tgt git aag gaa gaa gaa tgo 291 171 Pho Tyr Arg Asp Thr Val Ile Gly Asp Cys Val Lys Glu Glu Glu Cys 3 (1 173 dad caa cat gag and att cat gib tgaabgagag agbagtaata accaaaggito 346 174 App Gln His Glu Ile Ile His Val 45 176 maanttudge tolacaaaat egetagttigg atticteett tigogtigegaa tagtittaagt 406 178 tyuthttaay taasacutoo tgttgaayag aataaagett teeaaette 455 181 -010 - SEQ ID NO: 6 183 4211 · LENGTH: 75 183 -COLD - TYPE: PRT 184 GLIB · ORGANISM: Ascyclostoma caninum 186 (400) SEQUENCE: 6 188 Lys Ala Tyr Pro Glu Cys Gly Glu Asn Glu Trp Leu Asp Val Cys Gly Ç, 1.0 15 199 1 190 Thr Lys Lys Pro Cys Glu Ala Lys Cys Ser Glu Glu Glu Glu Asp

TIME: 13:22:09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/498,556C

Input Set : A:\Corvas-sequence listing.txt
Output Set: N:\CRF4\04152003\I498556C.raw

30 2.0 2.5 191 192 Pro Ile Cys Ang Ser Phe Ser Cys Pro Gly Pro Ala Ala Cys Val Cys 4 C 35 194 Glu Asp Gly Fhe Tyr Arg Asp Thr Val Ile Gly Asp Cys Val Lys Glu 5.0 -60 196 Glu Glu Cys Asp Glr. His Glu Ile Ile His Val 197 65 70 200 -010: SEQ ID NO: 7 201 - 211 - DENGTH: 81 mod of lmo TYPE: PET 1107 - 1117 OPGANISM: Ascyclostoma caninum DOS AGOOD SEQUENCE: 7 .07 Arg Thr Val Arg Lys Ala Tyr Pro Glu Cys Gly Glu Asr Glu Trp Leu 15 10 . 08 1 100 Asp Asp Cys Giy Thr Glr. Lys Pro Cys Glu Ala Lys Cys Asn Glu Glu .:10: 25 3.0 III Pro Fro Glu Glu Glu Asp Pro Ile Cys Arg Ser Arg Gly Cys Leu Leu . . $-4\,()$.17 Pm. Fro Ala Cys Val Cys Lys Asp Gly Pho Tyr Arg Asp Thr Val Ile ITS GLY Asp Cys Val Arg Glu Glu Glu Cys Asp Glr His Glu Ile Ile His 216 65 117 Val 110 -210 - SEQ 11 10: 8 7.11 +0.11 - LENGTH: 79 IL. - DII - TYPE: ERT JIA HILLA OF GAMISM: As cyclostoma caninum J. S. OROG - SEQUENCE: 8 [1] Ang Thi Mal Ang Lys Ala Tyn Pro Glu Cys Gly Glu Ash Glu Trp Leu 10 7 €. Lun Asp Val Cys Gly The Lys Lys Pro Cys Glu Ala Lys Cys Ser Glu Glu 230 .10 . Fl Glu Glu Glu Asp Pro Ile Cys Arg Ser Phe Ser Cys Pro Gly Pro Ala 3.5. 4 °. 4.0 ... Ala Tys Val Cys Giu Asp Gly Phe Tyn Arg Asp Thr Val Ile Gly Asp 234 50 · . Ę. -60135 Oys Val Lyc Gli Glu Glu Cys Asp Gln His Glu Ile Ile His Val 236 65 233 -210 - SEQ HD NO: 9 240 -011 - BENGTH: 711 .41 HLID - TYPE: DOM 24. (213 - ORGANISM: Andyclostoma deylanicum .44 -C200 - FEATURE: .45 -0.01 - NAME/KET: CDS $146 \times 21.1 \times \text{hodation:} (21) \dots (595)$ 24- KLZO - FEATURE: 249 -071 - NAME/KEY: MOD RES [950] CLOSE OTHER INFORMATION: Ecombinant cDNA Molecule AceNAP4 sequence L52 KALUZ SEQUENCE: 9

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/498,556C TIME: 13:22:09

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Gutput Set: N:\CRF4\04152003\I498556C.raw

2.54 gaattoacta ttatocaaca atq qoq qtq ott tat toa gta gca ata gog 50Met Ala Val Leu Tyr Ser Val Ala Ile Ala 11.5 1 5 157 tita ota otg gha toa caa tgo agt ggg aaa oog aac aat gtg atg act 98 ..! & Leu Leu Leu Val Ser Gln Cys Ser Gly Lys Pro Asn Asn Val Met Thr 1.4 15 760 ago got tut ggt out agt gas tat the get gag tgt ggd aat atg aag $146\,$.+1 Ash Ala Cys Gly Leu Ash Glu Tyr Phe Ala Glu Cys Gly Ash Met Lys 3.5 ${\mathbb N}({\mathbb R})$ qaa tigo quig cab ada tigo aat gag gag gaa aat gag gad agg gab gag ${\mathbb R} {\mathbb R}^4$ ightarrow4 Giu Cys Giu His Arg Cys Asn Giu Giu Giu Asn Glu Giu Arg Asp Giu 4.5 Led gam uga ata mon gow the etc ate out gtg that the out cot grat get 240 .x7 Glu Ard The Thr Awa Cys Leu Lie Arg Val Cys Phe Arg Pro Gly Ala 65 60 . 4.9 tigo qta tqo awa gwo gwa tto tat aga awo wga aca ggo ago tgt gtg 294 170 Cys Val Cys Lys Asp Gly Phe Tyr Arg Ash Arg Thr Gly Ser Cys Val 171 75 (4.1) TV, gas gas git gad two gag tab gag ast stg gag tto att sot ttt gds 33% .7% Glu Glu Amp Asp Cys Glu Tyr Gru Ash Met Glu Phe Ile Thr Phe Ala 10% doa gaa gha bog ata tgt ggt tod aad gaa agg tad tod gad tgd ggd (306 276 Pro Glu Val Pro Ile Cys Gly Ser Ash Glu Arg Tyr Ser Asp Cys Gly 110 115 JTM aut que que cau cas toe gay ege aus tye auc gay gae gat tat gay aug 434 200 Ash Asp Dys Gln Cys Glu Arg Dys Cys Ash Glu Asp Asp Tyr Glu Dys 130 1.351.15 Lef gga gat gag goa two ogo toa oat gtt tot gaa ogt oot ggt goo tot 450 . F. Gly Asp Glu Ala Cys Ang Sen His Val Cys Glu Ang Pro Gly Ala Cys 145 UPA qua ngo gwa gwo ggg tto tao aga aac aaa asa ggt ago tgt gtg gaa 550 184 Val Cys Gli Asp Gly Phe Tyr Ang Ash Lys Lys Gly Sen Cys Val Glu 286 155 160165 .987 ago gat gad tgo gwa tao gat aat atg gat tto ato abt tit goa coa 578 THE Ser Asp Asp Cys Glu Tyr Asp Ash Met Asp Phe Ile Thr Phe Ala Pro 175 185 160 mun qaa abb toa oga taaccasaga tgctacotot ogtacgoaac tccgctgatt gaggtt 636, 741 Glu Thr Ser Arg 200 1.00190 gauthanten etigeakete aacattitit tigigatgot gigeatetga gettaacetg 696 711 195 ataaagoota tggtg. 298 -210 - SEQ IE NO: 10 144 - 1211 - BENGTH: 425 300 HL12 - TYPE: DNA 301 -0013 - ORGANISM: Amoyolostoma ceylanicum Sub-COO-FEATURE: FOR HELLI - NAME/KEY: CDS $\pm 6.5 \pm 0.322 \pm 100 \text{ATION:} (10) \dots (291)$ 30 ' -1 20 / FEATURE:

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/498,556C

DATE: 04/15/2003 TIME: 13:22:10

Input Set: A:\Corvas-sequence listing.txt
Cutput Set: N:\CRF4\04152003\I498556C.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

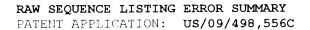
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Seq#:130; Kaa Fos. ., /, 4, 5, 6
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/498,556C

DATE: 04/15/2003 TIME: 13:20:10

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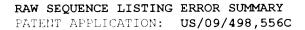
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Ceq#:165; Maa Fes. 1,.,e,5,6,7,5,9,10,11,1.,13,14,15,10,17,18
Neq#:166; Maa Fess. 1,7,8,5,6,7,-,9,10,11,12,13,14,15,16,17
Seq#:167; Maa Pos. 1,.,3,5,6,7,8,9,10,11,1_,13,14,15,16
Seq#:168; Haa Pus. 1, ., 5, 5, 6, 7, 8, 9, 10, 11, 15, 13, 14, 15
Thig#:169; Maa P.s. 1,.,-,5,6,7,8,9,10,11,10,13,14
So g#:170; Maa Eus. 1, 3, 3, 5, 8, 7, 4, 4, 10, 11, 12, 13
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C-q#:178; Maa Pos. 3,3,4
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Fog#:188; Maa Fos. 7, 3,4,5,6
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Seq#: 227; Maa Fos. ., 3, 4
Seg#:2.8; Maa Pos. 1,4,5,6,7,8,4,10,11,12,13,14,15
Seq#:219; Maa Fos. 1,4,5,6,7,8,9,10,11,12,13,14
Seq#:230; Maa Fos. 2,3,4,5,6,7,8
Seq#:331; Kaa Fos. 3,:,4,5,6,7
Seq#:032; Maa Pos. 2,3,4,5,6
Seq#:133; Maa Fos. 7,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22
Seq#:133; Maa Pos. U3,04,15,26
Seq#:134; Maa Fos. 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 11, 22
Seq#:234; Maa Pos. 13,24,25
Seq#:235; Maa Pos. 0,5,4,5,6,7,5,9,10,11,10,13,14,15,16,17,18,19,00,01,00
Seq#:235; Haa Pos. .3,24
Seq#:236; Maa Fos. 1,3,4,5,6,7,8,9,10,11,10,13,14,15,16,17,18,19,00,21,00
Seq#:236; Maa Pos. 33
Seq#:237; Maa Pos. 0.3, 3.4, 5.6, 7.8, 9.10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22
Seq#:236; Maa Pos. 2, 4,4,5,6,7,8,9,10,11,1.,13,14,15,16,17,18,19,30,21
Seq#:239; Maa Pos. 1, %, 4, 5, 6, 7, %, 9, 10, 11, 1., 15, 14, 15, 16, 17, 18, 19, 20
Steq#::340; Maa Pos. .,3,4,5,6,7,8,0,10,11,15,13,14,15,16,17,16,19
Seq#::341; Maa Pos. ., 3, 4, 5, 6, 7, 8, 9, 10, 11, 10, 13, 14, 15, 16, 17, 18
Seg#: 240; Maa Pos. 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17
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DATE: 04/15/2003 TIME: 13:22:10

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Cutput Set: N:\CRF4\04152003\I498556C.raw

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Seg#:244; Maa Fos. 1,3,4,5,6,7,8,9,10,11,10,13,14,15
Seq#:.45; Maa Fos. 2,0,4,5,6,7,8,0,10,11,10,13,14
Seq#:140; Maa Fos. 1,0,4,5,6,7,8,3,10,11,10,13
Seq#:.4"; Maa Fos. .,0,4,5,+,7,6,+,10,11,10
Seq#: 48; Maa Fos. ., 3, 4, 5, 6, 7, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}
Seq#:.44; Kaa Ecs. .,?,4,5,+,7,8,5,10
Seq#::50; Maa Fcs. 1,2,4,5,6,7,5,4
Seq#: 51; Maa Pos. 0,0,4,6,6,6,7,8
Seq#: 25%; Maa Fos. 2,8,4,6,6,7
Seq#:.53; Kaa Fos. 2,3,4,5,6
Seig#:114; Maa Fos. 1,2,4,5,6,7,8
Seq#: 155; Maa Fos. 1, 2, 4, 5, 6
Seq#:250; Maa Pos. 2,2,4,5
Seq#: 57; Maa Pos. 1,0,4
Seq#: Se; Naa Pes. 1,1,4,6,6,7,8,4,10,11,12,13,14,15,16,17,18,19,20,21,22
Seq#: 58; Maa Fos. 13
Seq#: 50; Kaa Fos. 1,0,-,-,0,7,8,9,10,11,10,13,14,15,16,17,18,19,10,01,00
Seq#:: 60; Maa Fos. 1, 2, 5, 6, 6, 7, 8, 9, 10, 11, 13, 13, 14, 15, 16, 17, 18, 19, 10, 31
Seq#:..61; Maa Pos. 1,.,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,10
Seq#: Not; Haa Fos. 1, 1, 3, 5, 4, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19
Seq#::00; Maa Fes. 1, 2, 3, 6, 6, 7, 3, 9, 10, 11, 13, 13, 14, 15, 16, 17, 18
Seq#::de4; Maa Fos. 1,1,7,8,6,6,7,8,9,10,11,11,13,14,15,16,17
Seq#:: 66; Maa Fos. 1,2,5,6,6,7,6,9,10,11,12,13,14,15,16
Seq#:: 66; Maa Fos. 1,2,5,6,6,7,6,6,10,11,12,13,14,15
Seq#: (6); Maa Pos. 1, ., ., 1, 6, 7, 6, 3, 10, 11, 1., 13, 14
Seq#: 66; Maa Fes. 1,2,:,0,7,6,3,10,11,10,13
Seq#: 69; Maa Pos. 1,1, 1,1,6,7,8,9,10,11,1.
Seq#:.70; Maa Pos. 1,2,4,8,6,7,8,9,10,11
Seq#:171; Maa Fos. 1,2,3,5,6,7,5,9,10
Seq#: 70; Kaa Pos. 2,7,4,5
Seq#: 173; Maa Pos. 1, 2,4
Seg#: /4; Maa Pos. 2,3,4,5,6
Seq#:275; Maa Pos. 0,:,4,5
Seq#:176; Naa Pos. 2,:,4
Seq#: 77; Maa Pos. .,4,5,6,7,8,9,10,11,12,13,14,15
Seq#:17-; Maa Pos. 1,4,5,6,7,8,9,10,11,12,13,14
Seq#:::/H; Maa Pos. .,4,5,6,7,8,9,10,11,12,13
Seq#: 380; Kaa Pcs. 1, 1, 4, 5, 6, 7, 8
Seg#:::81; Maa Pos. 2,5,4,5,6,7
Seq#: Maa Pos. 1,2,4,5,6
Seq#:283; Kaa Fos. 2,:,4,5,6,7,8,9,10,11,10,13,14,15,16,17,13,19,20,21,22
Seq#:::8x; Maa Pos. .3,24,25,26
Seq#: 184; Maa Pos. 13,24,25
Seq#:. M°; Maa Fos. 1, 5,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,30,31,33
Seg#: 185; Maa Pos. 13,24
Seq#::M6; Maa Pes. ., 5, 4, 1, 6, 7, 6, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22
Seg#:28n; Maa Pos. 13
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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/498,556C

DATE: 04/15/2003 TIME: 13:22:10

Input Set : A:\Corvas-sequence listing.txt
Ou.put Set: N:\CRF4\04152003\I498556C.raw

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Seq#:287; Maa Fos. 2,8,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22
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Seq#:::89; Maa Pos. :::,0,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,::0
Seq#:290; Maa Fos. 2,5,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
Seq#:290; Maa Fos. 2,3,4,5,6,7,8,9,10,11,10,13,14,15,16,17,18
Seq#:290; Maa Fes. 0,3,4,5,6,7,8,9,10,11,10,13,14,15,16,17
Seq#:::190; Maa P. S. E., S., 4, 5, +, 7, F., 4, 10, 11, 12, 13, 14, 15, 16
Sug#::394; Maa Fes. 2,8,4,5,6,7,8,9,10,11,12,13,14,15
Seq#:295; Maa Fos. 3,3,4,5,6,7,8,9,10,11,13,13,14
Seq#:236; Maa Fes. 3,5,4,5,6,7,-,9,10,11,13,13
Seq#::::97; Maa Fes. 3,3,4,5,4,7,5,3,10,11,13
Seq#:290; Maa F s. 3,5,4,5,6,7,6,9,10,11
Seq#: 149; Maa Fos. 3,3,4,5,6,7,8,4,10
Seq#:300; Maa Fos. 3,3,4,5,6,7,5,4
Seq#: 001; Maa Pos. 1,0,4,5,6,7,8
Seq#: 200; Maa Prs. 1,3,4,5,6,7
Ser#::03; Maa Pos. 1,2,4,5,6
Seq#:304; Maa Ens. 1,5,4,5
Sleg#: 805; Maa Evs. 1,5,4
Sug#: 30h; Kaa Eus. .,
Seg#:::07; Maa Fes. .
Seq#:308; Maa Fos. .,3,4,5,6,7,8
Seq#:309; Maa Frs. 2,3,4,5,6
Seq#:310; Maa Fos. .,5,4,5
Seg#: 511; Maa Ers. ., 1,4
Seq#::10; Maa Fes. 1,1,7,5,6,7,8,9,10,11,10,13,14,15,16,17,18,19,20,01,22
Seg#: Bld; Maa Fes. .3
Seq#:315; Maa Fos. 1,5,5,6,7,5,4,10,11,10,13,14,15,16,17,18,19,00,01,22
Seq#:314; Haa Frs. 1,2,3,5,6,7,8,4,10,11,12,13,14,15,16,17,18,19,20,21
Stood#: 315; Maa Pos. 1,7,2,5,6,7,3,4,10,11,12,13,13,15,14,15,16,17,18,19,20
Seq#::110; Maa Fes. 4,0,2,5,0,7,8,9,10,11,10,10,10,14,15,10,17,18,19
Seq#::1/; Maa Fes. F,.,F,S,C,7,5,0,10,11,12,15,14,15,16,17,18
Seq#: 41%; Maa Fos. 1,2,3,5,4,7,5,3,10,11,13,13,14,15,16,17
Beq#:319; Maa Fos. 1,.,5,5,6,7,4,4,10,11,13,15,14,15,10
Seq#:300; Maa Fos. 1,2,1,5,6,7,8,4,10,11,10,13,14,15
Seq#: 331; Maa F. s. 1,.,3,5,0,7,8,9,10,11,13,15,14
Geq#:Phh; Maa Fus. 1,2,2,5,6,4,7,8,9,10,11,10,13
Seq#:528; Maa Fes. 1,5,5,6,7,8,9,10,11,12
Seq#: 534; Maa Fos. 1,., 5,8,10,7,8,9,10,11
Sog#:500; Maa Fos. 1,2,2,5,6,7,8,9,10
Seig#: 326; Maa Frs. 2, 2, 4, 5
Seg#: 3.17; Maa Pos. 1, 1,4
Seq#: 328; Maa Pos. 2, 1, 4, 5, 6
Seq#:309; Kaa Prs. 2,:,4,5
Seig#:330; Maa Pus. 3,3,4
Seg#: \mathbb{R}^{2}; Maa I s. \mathbb{S}, 4, 5, 6, 7, 8, 4, 10, 11, 12, 13, 14, 15
Seq#: vid; Maa Fis. 3,4,5,6,7,8,9,10,11,12,13,14
Seq#: 3:3; Maa Fos. 3, 4, 4, 5, 6, 7, 8
Seg#: 534; Xaa Los. 2,5,4,5,6,7
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/15/2003 FATENT APPLICATION: US/09/498,556C TIME: 13:13:10

Input Set: A:\Corvas-sequence listing.txt
Output Set: N:\CRF4\04152003\I498556C.raw

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Seq#:335; Kaa Pos. 1,3,4,5,6
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Seq#:33%; Maa Pos. 13,24,25,26
Seq#:33%; Maa Fos. L,3,4,5,+,7,8,+,10,11,1L,13,14,15,16,17,18,19,00,01,00
Seq#:337; Kaa Fcs. 23,34,25
Seq#:33%; Maa Fos. 7,3,4,5,6,7,8,3,10,11,1.,13,14,15,16,17,18,19,00,01,00
Seq#:33F; Maa Fos. 13,24
Seq#:339; Maa Fos. 2,3,4,5,4,7,8,9,10,11,1.,13,14,15,16,17,18,19,00,01,00
Seq#:33 : Maa Pos. .3
Seq#:340; Maa Pos. 1,3,4,5,6,7,8,9,10,11,10,13,14,15,16,17,18,19,00,01,02
Seq#:341; Maa Fos. 2,3,4,8,6,7,8,4,10,11,10,13,14,15,16,17,18,19,00,01
Seq#:341; Maa Pos. 2,3,4,5,6,7,6,4,10,11,10,16,14,15,16,17,16,19,20
Seq#:340; Maa Fos. 1,3,4,5,6,7,6,9,10,11,12,13,14,15,16,17,18,19
Seq#:344; Maa Fos. 1,3,4,5,+,7,8,9,10,11,12,13,14,15,16,17,18
Seq#:345; Maa Pis. .,3,4,5,+,7,5,4,10,11,15,13,14,15,16,17
Seq#:340; Maa Fos. 1,3,4,5,7,3,4,10,11,11,13,13,14,15,16
Seq#:347; Maa Fos. .,3,4,8,+,7,3,4,10,11,1.,13,14,15
Seq#:34%; Maa Fis. 2,3,4,5,6,7,8,9,10,11,10,13,14
Seq#:34+; Maa Ecs. L,3,4,5,6,7,9,9,10,11,10,13
Seq#:350; Maa Fos. 2,3,4,5,6,7,8,4,10,11,12
Seq#:351; Naa Fos. 2,3,4,5,6,7,6,4,10,11
Seq#:350; Maa Pos. 1,3,4,5,6,7,8,4,10
Seq#:353; Maa Pos. 2,3,4,5,6,7,8,9
Seq#:354; Maa Pos. 1,3,4,5,6,7,6
Seq#:355; Maa Pos. 0,3,4,5,6,7
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Seq#:357; Maa Pos. .,4
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:68; Line(s) 1996,2001